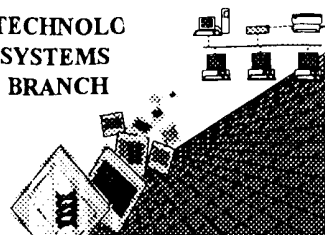


5040

BIOTECHNOLOGY
SYSTEMS
BRANCH



RAW SEQUENCE LISTING
ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:

10/030,658

Source:

P4/10

Date Processed by STIC:

2/8/2002

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FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

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Revised 01/29/2002



PCT10

RAW SEQUENCE LISTING

DATE: 02/08/2002

PATENT APPLICATION: US/10/030,658

TIME: 10:31:52

Input Set : A:\PTO.VSK.txt

Output Set: N:\CRF3\02082002\J030658.raw

Does Not Comply
Corrected Diskette Needed

3 <110> APPLICANT: Yamamura Ken-ichi
4 Araki Kimi
6 <120> TITLE OF INVENTION: TRAPVECTOR
8 <130> FILE REFERENCE: PH-976PCT
10 <140> CURRENT APPLICATION NUMBER: US/10/030,658
10 <141> CURRENT FILING DATE: 2002-01-11
10 <150> PRIOR APPLICATION NUMBER: JP99/200997
11 <151> PRIOR FILING DATE: 1999-07-14
E--> 13 <160> NUMBER OF SEQ ID NOS: (12/14) (p2)
15 <170> SOFTWARE: PatentIn Ver. 2.0

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128 <211> LENGTH: 400
129 <212> TYPE: DNA
130 <213> ORGANISM: Mus musculus
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142 <211> LENGTH: 416
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Input Set : A:\PTO.VSK.txt
Output Set: N:\CRF3\02082002\J030658.raw

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181 gatgcccag gggataagct ggagctgtca g 211
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Item 9

last sequence in file

delete

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/030,658

DATE: 02/08/2002

TIME: 10:31:53

Input Set : A:\PTO.VSK.txt

Output Set: N:\CRF3\02082002\J030658.raw

L:10 M:270 C: Current Application Number differs, Replaced Current Application No
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:133 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:11
M:340 Repeated in SeqNo=11
L:147 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:12
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L:161 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:13
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L:177 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:14 differs:12
L:184 M:254 E: No. of Bases conflict, LENGTH:Input:1 Counted:212 SEQ:14
L:184 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:184 M:252 E: No. of Seq. differs, <211>LENGTH:Input:211 Found:212 SEQ:14
L:13 M:203 E: No. of Seq. differs, <160> Number Of Sequences:Input (12) Counted (14)